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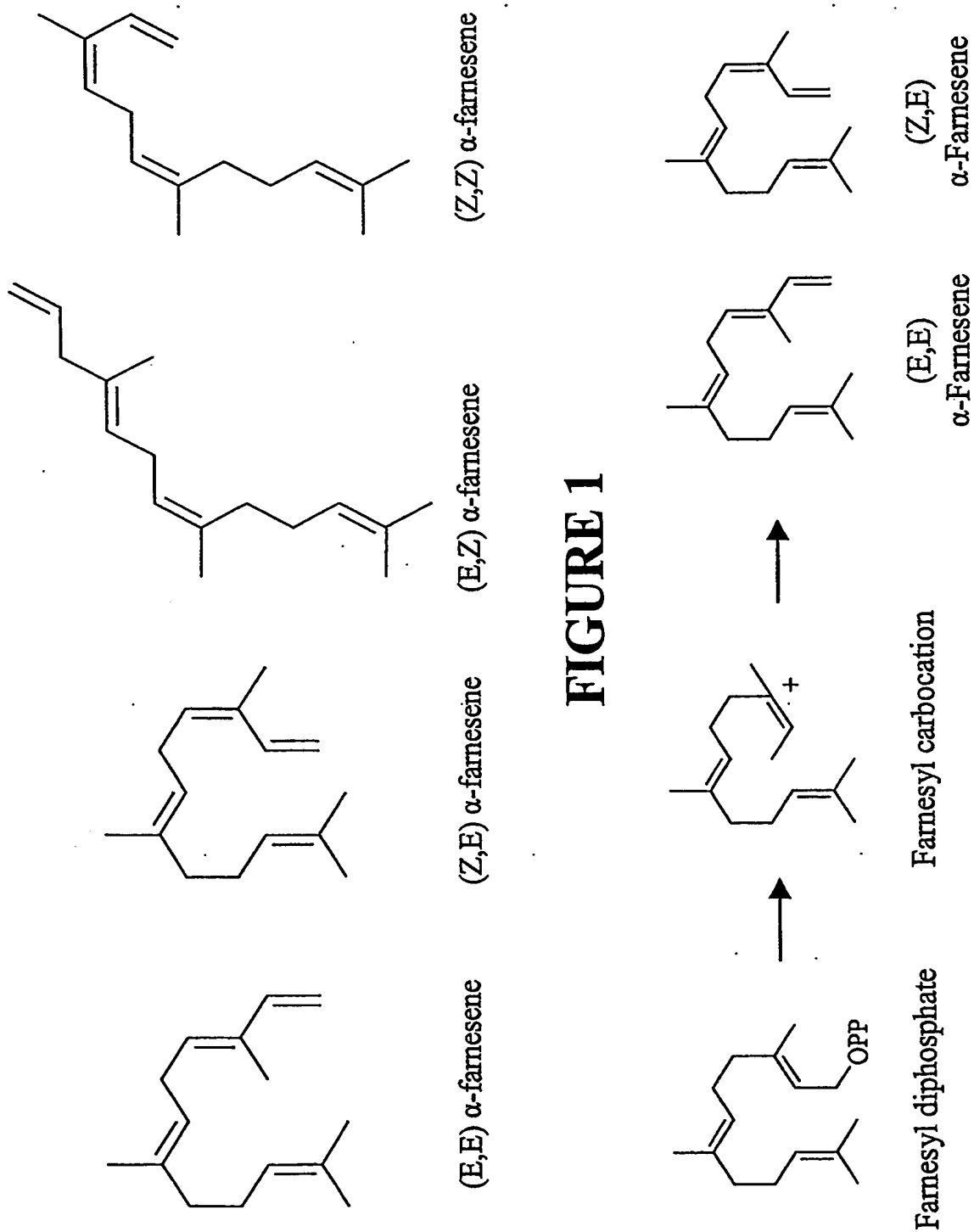


FIGURE 1

FIGURE 2

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1 CTATAGCTTC TTGTAATCCA AACATCTCGA GCTTCTTGTA CACCAATTA GGTATTCACT ATGGAATTCA GAGTTCACCT GCAAGCTGAT AATGACGAGA
101 AATATTTTCA AAACCAAGATG AAACCCGAAC CTGAAGCCTC TTACTTGATT AATCAAGAC GGTCTGCCAAA TTACAGCCA AATATTTGGA AGAAGCAATT
201 CCTAGATCAA TCTCTTATCA GCAATATCGA TGGAGATCAG TATCGGAAGC TGCTTCAGAA GTTATAGAA GAAGTTAAGA TTTATATATC TGCTGAACA
301 ATGCAATTAG TAGCTAAGTT GGAGCTCATT CACAGGCTCC GAAAAGTAGG CCTCGGGAAC CTCCTGCAAA AGGAATCAA GGAAGCCCTA GACAGCAATTG
401 CAGCTATCGA AAGCGACAAT CTCGGGACAA GAGACGATCT CTATGGTACT GCATTACACT TCAAGATCCT CAGGCAGCAT GGCTATAAAG TTTCACAAGA
501 TATATTGGT AGATTTCATGG ATGAAAAGGG CACATTAGAG AACCAACATT TCGCGGATTT AAAAGCAATG CTGGAACCTT TCGAGCCCTC AAACCTGGGT
601 TTCGAGGGTG AAGATATTTT AGATGAGGGG AAGCTTCTCT TGACGCTAGC TCTCAGAGAT AGTGGTCATA TTTCTTATCC AGACAGTAAC CTTTCCAGGG
701 ACGTAGTTCA TTCCCTGGAG CTTCCATCAC ACCGCAGAGT GCAGTGGTTT GATGTCAAAI GCGAATCAA CCGCTATGAA AAAGACATTT GTCGGCTCAA
801 CCCCACCTTA CTCGAATTAG CAAGCTTAA TTTCAAGCTA GTTCAGGCCC AACTCCAAA AAACCTAAGG GAAGCATCCA GGTGGTGGCG AAATCTGGCG
901 TTGGCAGACA ACTTGAAATT TGCAAGAGAT AGACTGGTTG AATGTTTCTC ATGTGCTGTG GGACTAGCAT TCGAGGCTGA GCACTCATCT TTTAGAATAT
1001 GTCTTACCAA AGTCATCAAC TTAGTACTGA TCATACAGCA GGTCTATGAT ATTTATGGCT CAGAGGARGA GCTAAGGCAC TTCACCAATG CTGTTGATAG
1101 CTGGGATTCT AGGGAACCTG ACCAGCTTCC AGAGTGTATG AAGATCTGTT TCCAAGTACT CTACAACACT ACTTGTGAAA TTGCTCGTGA AATTGAGGAG
1201 GAGAAATGCT GGAACCAAGT ATTACCTCAA TTGACCAAG TCTGCCGAGA TTTTGTAAA GCATTATTGG TCGAGGCGAGA GTGCTATATAT AAGAGCCATA
1301 TACCAACCCCT TGAAGACTAC CTAAGAAAGG GATGCATTTT ATCATCAGTT TCAGTGTCTT TCAGTTCACCT GTTTTCTCT ATAACTCATG AGGGAACCAA
1401 AGAGATGGCT GATTTCTTTC ACAAGAATGA AGATCTTTTG TATAATATCT CTCTCATCGT TCGCCTCAAC AATGATTTGG GAACCTCCGC GGCTCAACAA
1501 GAGAGAGGGG ATTCCTCTTC ATCAATCGTA TGTTACATGA GAGAAGTGA TCCCTCTGAA GAACAGCTA GGAAGAACAT TAAGGCCATG ATAGACAATG
1601 CATGGAAGAA ACTAATGGA AATGCTTCA CAACAACCA AGTGCCTTTT CTGTGATCAT TCATGAACAA TGCCACAAC ATGCCACCTG TCGCCACACAG
1701 CCTTTACAAA GATGGAGATG GGTTTCCGTGA CCAAGAGAAA GGGCCCTCGA CCGCATCTCT GTCCTTACTA TTCCAACCTC TTGTAACCTA GTACTCATAT
1801 ACTTTGAAAT AAATAGCAGC AAGAACTTTG CGGTTGAGTT CGTCAATGAT AAATTAATCT TTACAGTTTG TAACCTTGT GCACAAGAT TATGAATAAA
1901 AAGTTCTAGT TTGCTGTTA TTTTAAAAA AAAAAAAA AAAAAAAA AA

FIGURE 3

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1 MEFRVHLQAD NEQKIFQNQM KPEPEASYLI NQRRSANYKP NIWKNDFLDQ
50 SLISKYDGDE YRKLSEKLE EVKIYISAET MDLVAKLELI DSVRKLGLAN
100 LFEKEIKEAL DSIAAIESDN LGTRDDLYGT ALHFKILRQH GYKVSQDIFG
150 RFMDEKGTLE NHHFAHLKGM LELFEASNLG FEGEDILDEA KASLTALRD
200 SGHICYPDSN LSRDVVHSLE LPSHRRVQWF DVKWQINAYE KDICRVNATL
250 LELAKLNFNV VQAQLQKNLR EASRWWANLG FADNLKFARD RLVECFSCAV
300 GVAFEPEHSS FRICLTKVIN LVLIIDDVYD IYGSEEELKH FTNAVDRWDS
350 RETEQLPECM KMCQVLYNT TCEIAREIEE ENGWNQVLPQ LTKVWADFCK
400 ALLVEAEWYN KSHIPTLEEY LRNGCISSSV SVLLVHSFFS ITHEGTKEMA
450 DFLHKNEDLL YNISLIVRLN NDLGTSAAEQ ERGDSPSSIV CYMREVNASE
500 ETARKNIKGM IDNAWKVNG KCFTTNQVPF LSSFMNNATN MARVAHSLYK
550 DGDGFGDQEK GPRTHILSLL FOPLVN*

FIGURE 4

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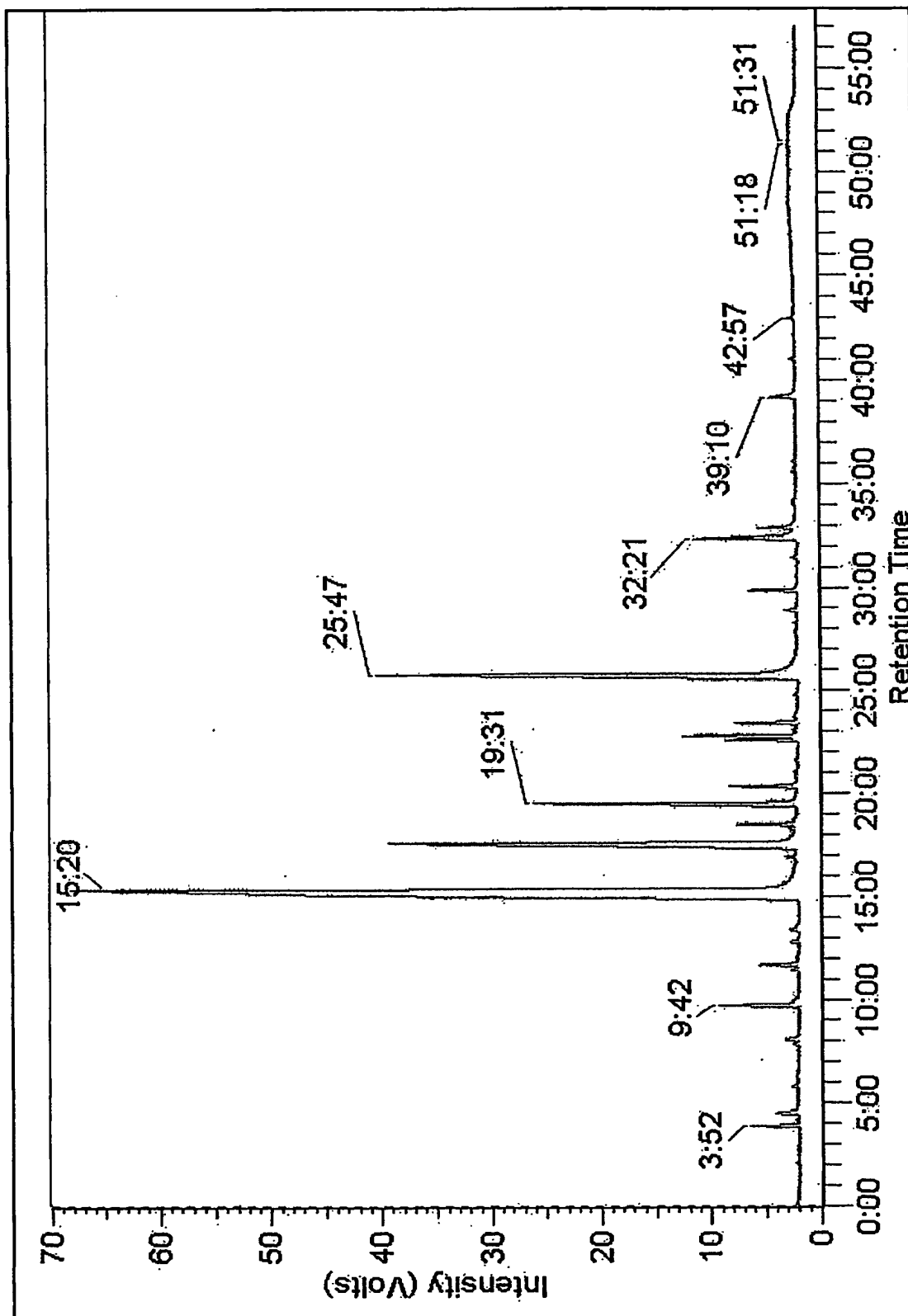


FIGURE 5

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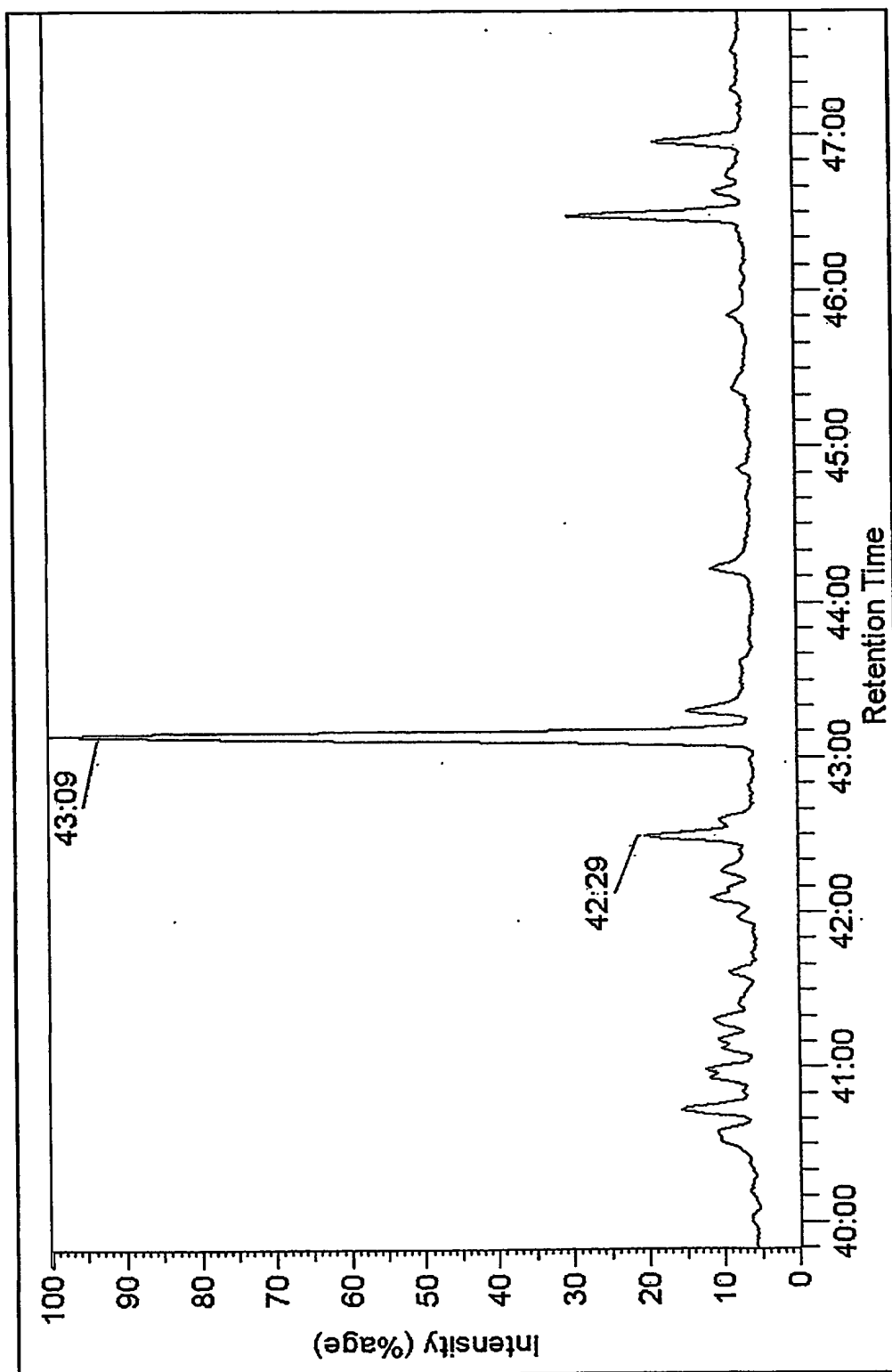
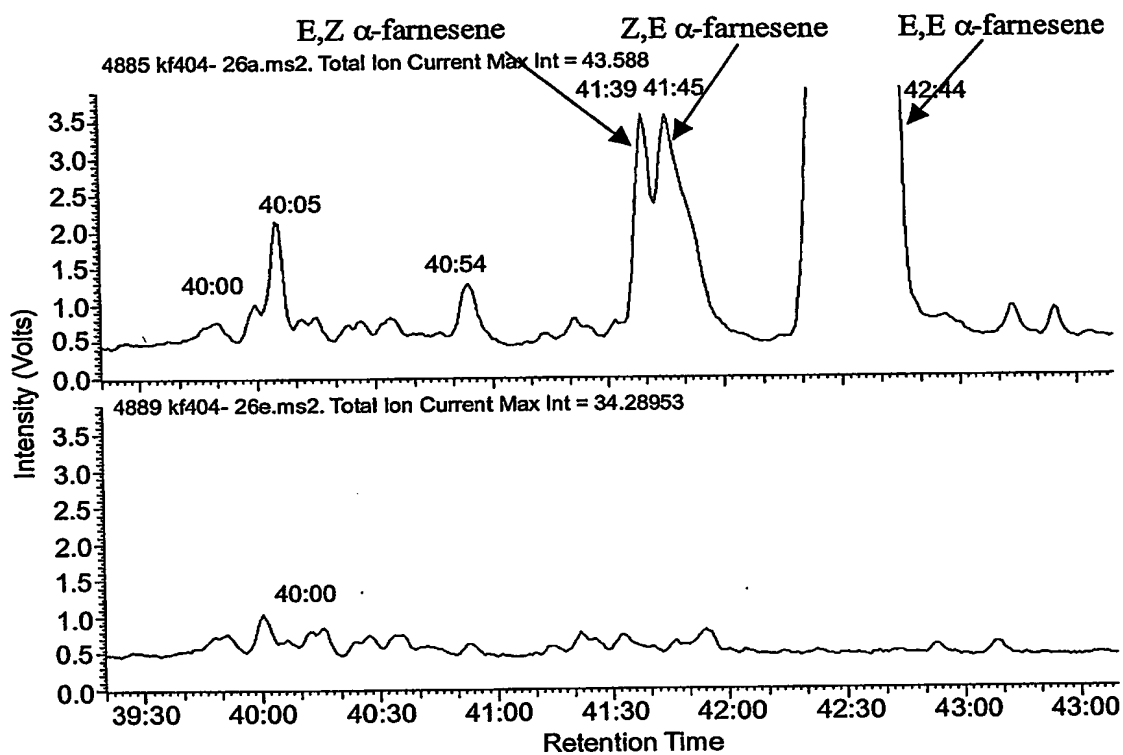
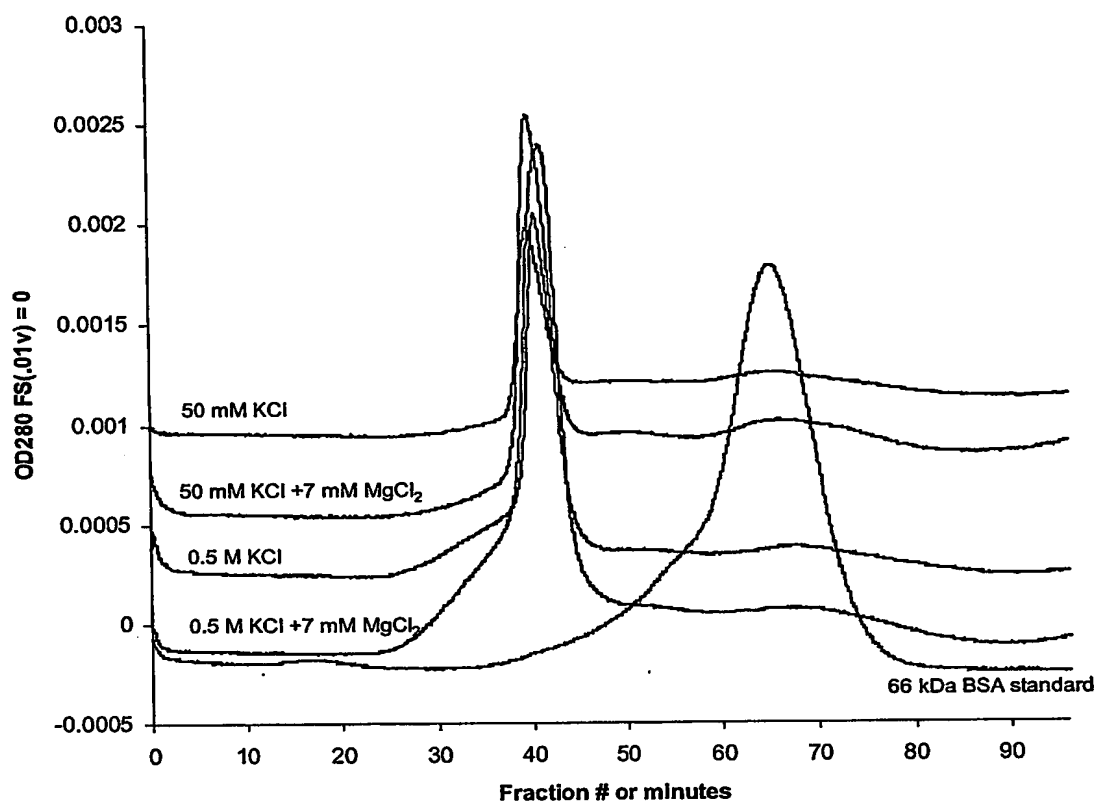


FIGURE 6

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**FIGURE 7**

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**FIGURE 8**

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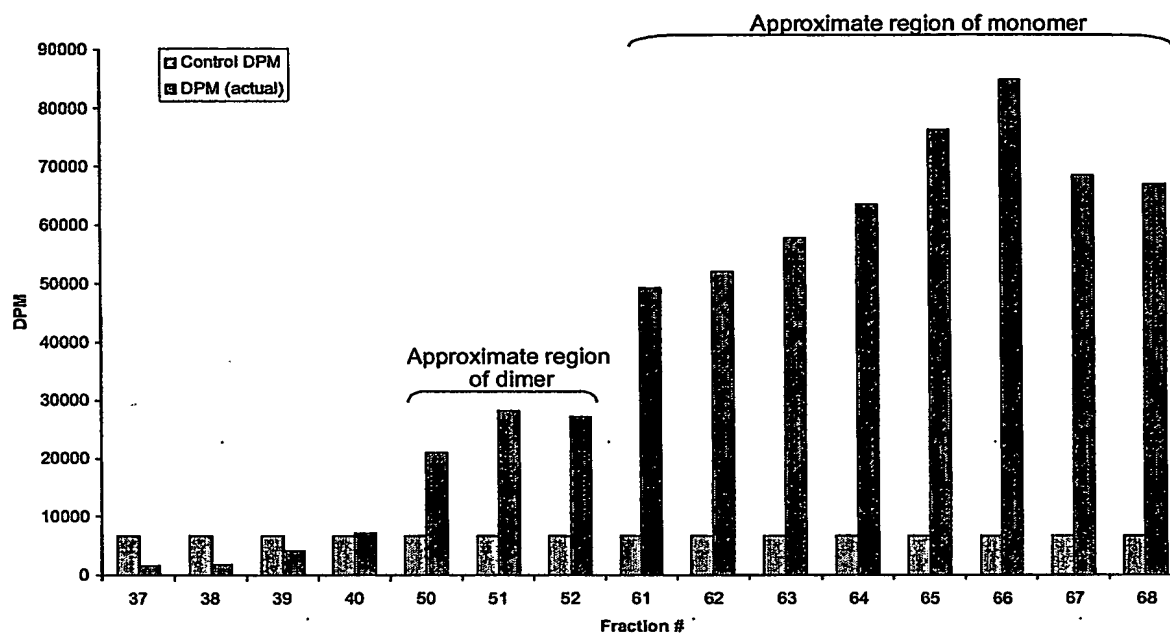


FIGURE 9

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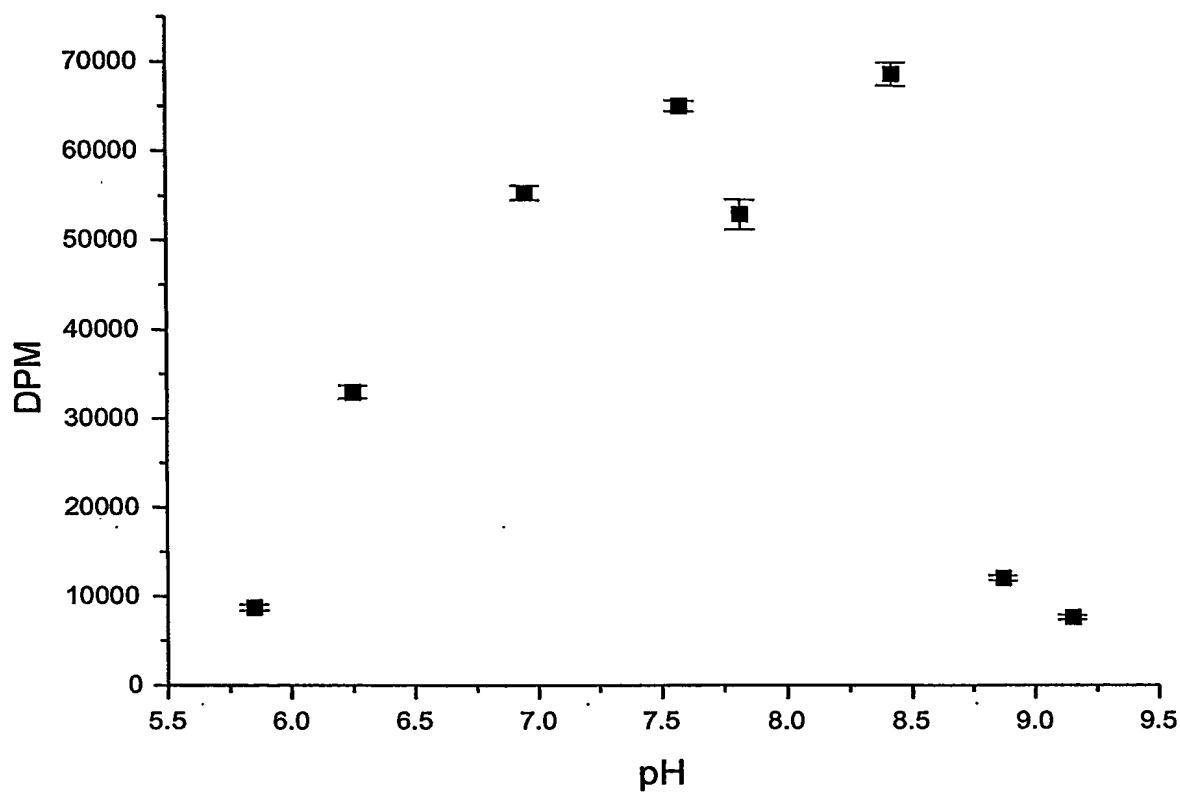
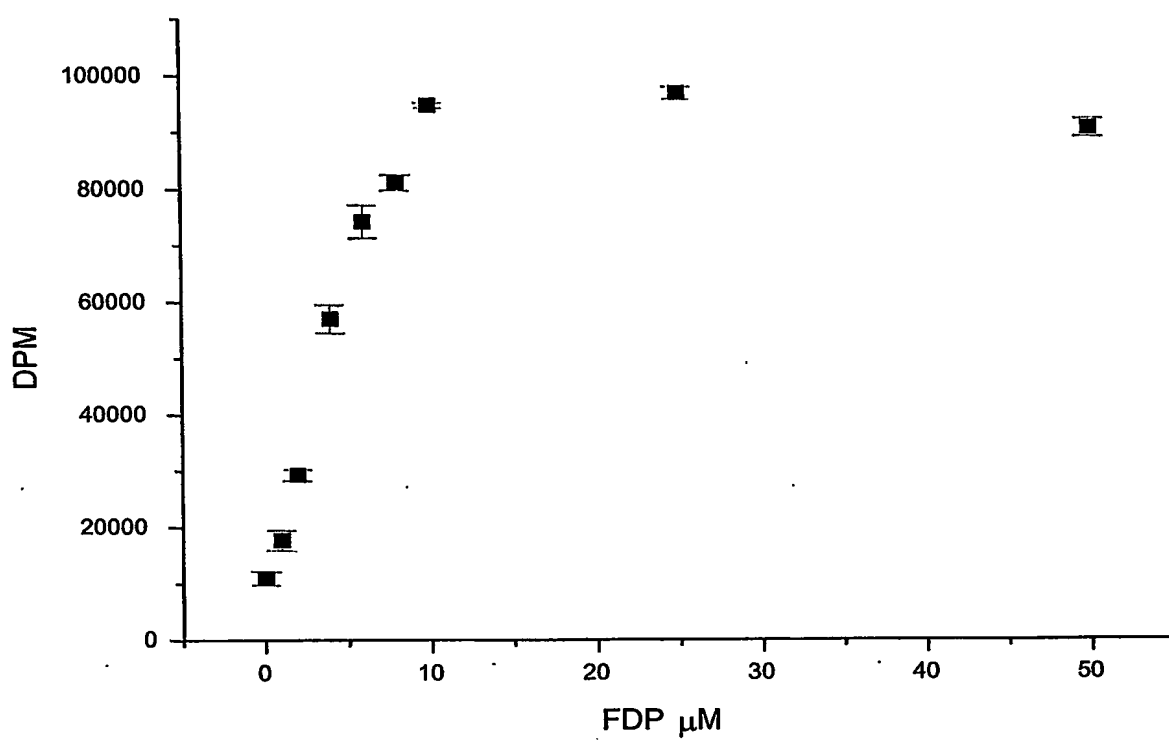
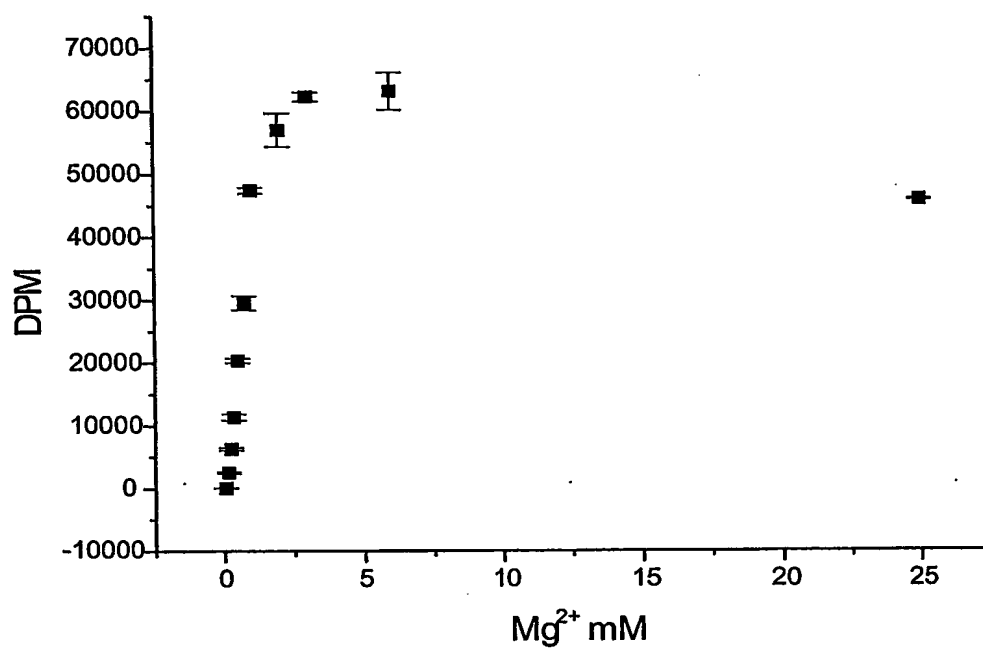


FIGURE 10

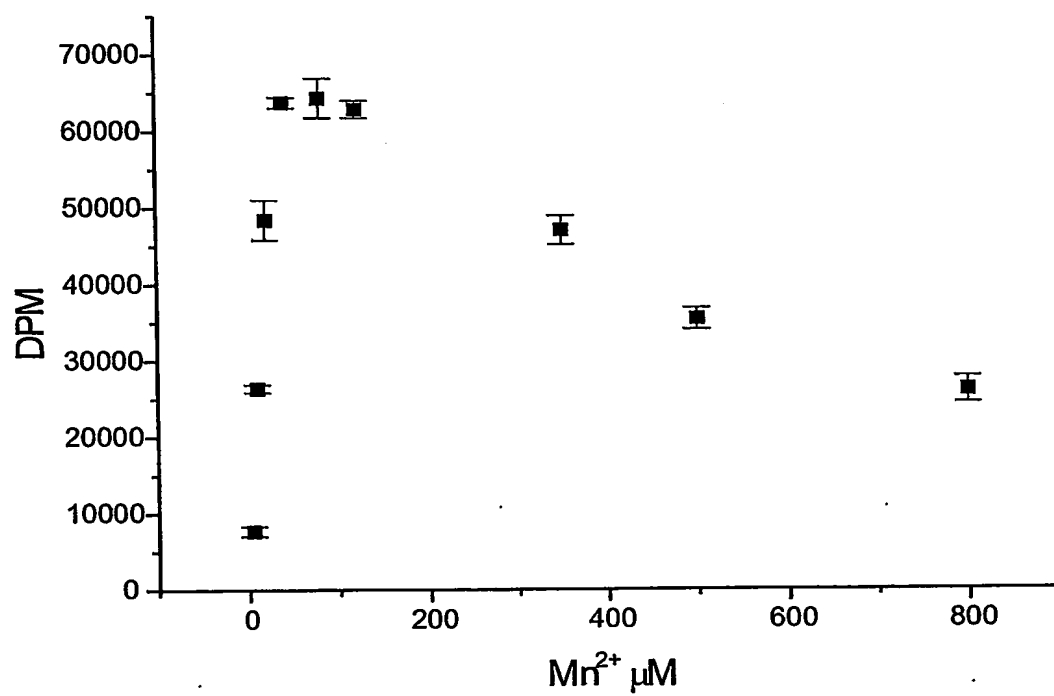
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**FIGURE 11**

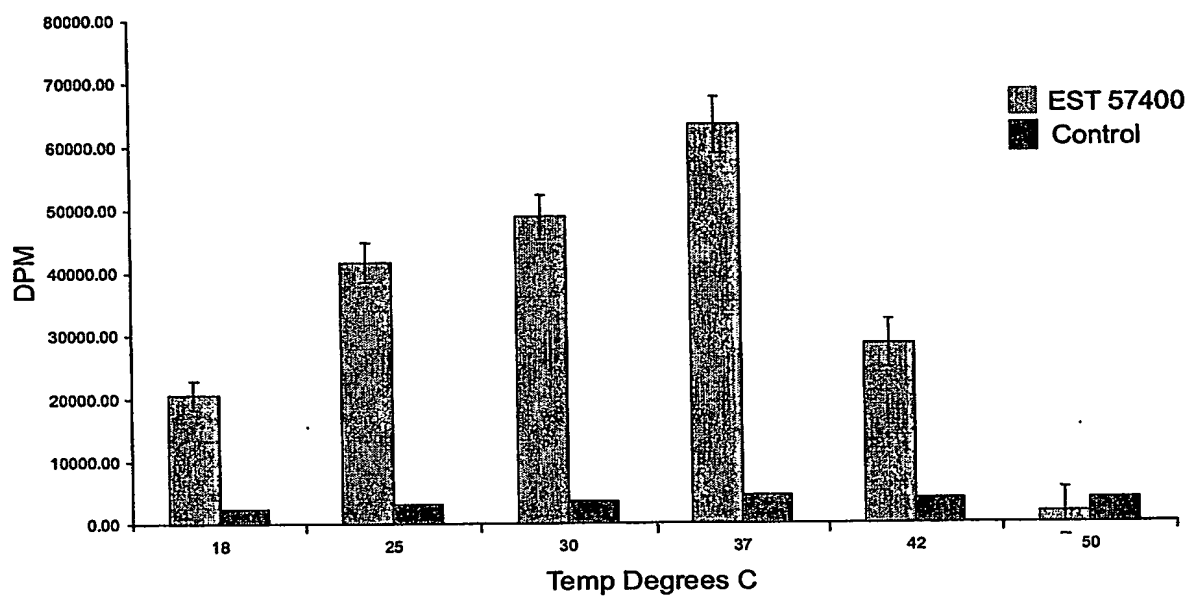
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**FIGURE 12**

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**FIGURE 13**

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**FIGURE 14**

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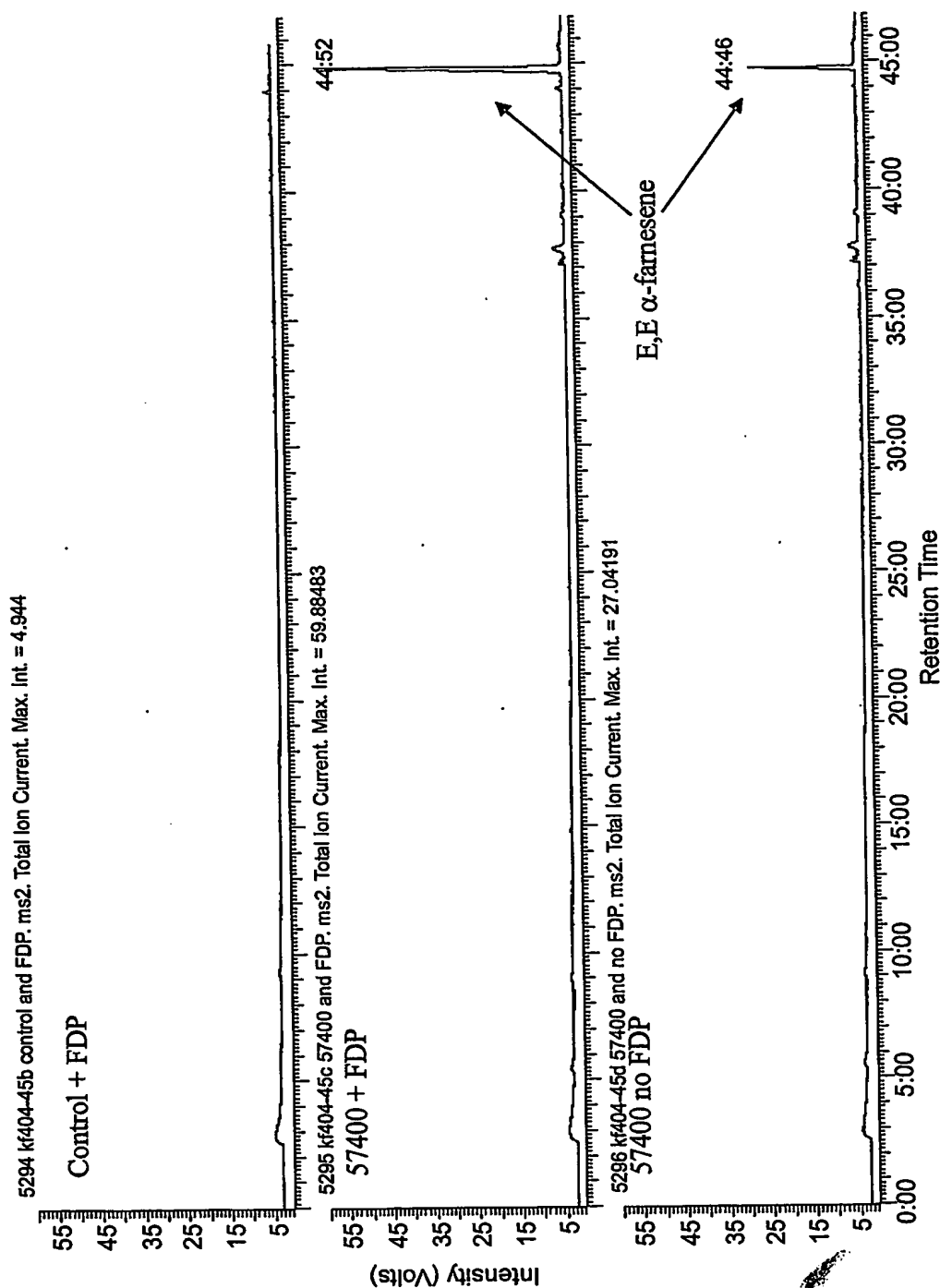


FIGURE 15

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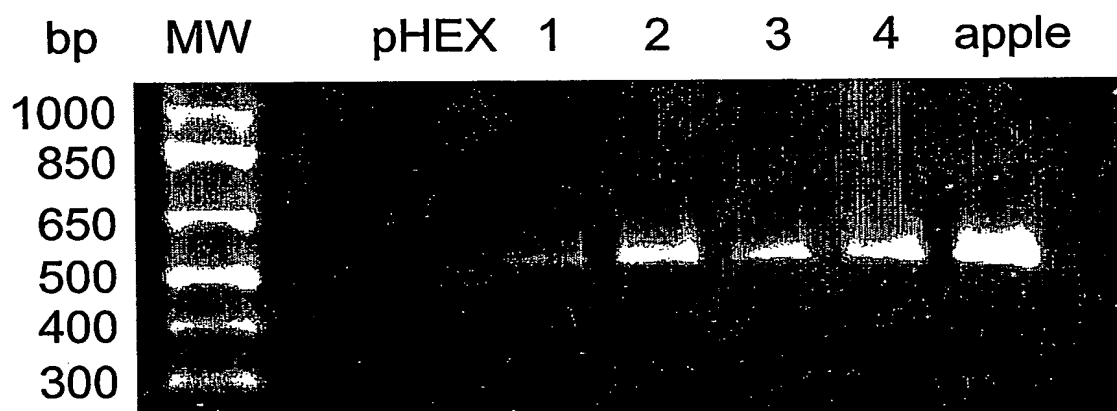


FIGURE 16

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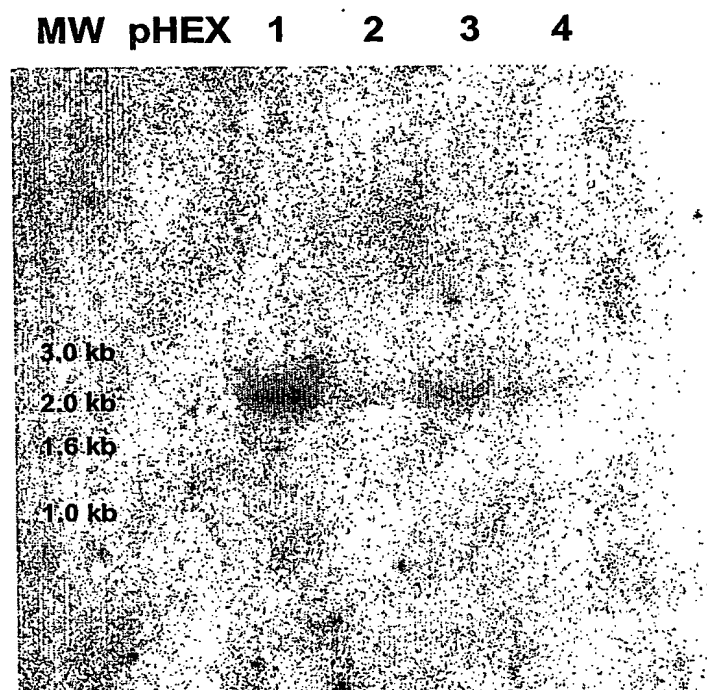


FIGURE 17

FIGURE 18

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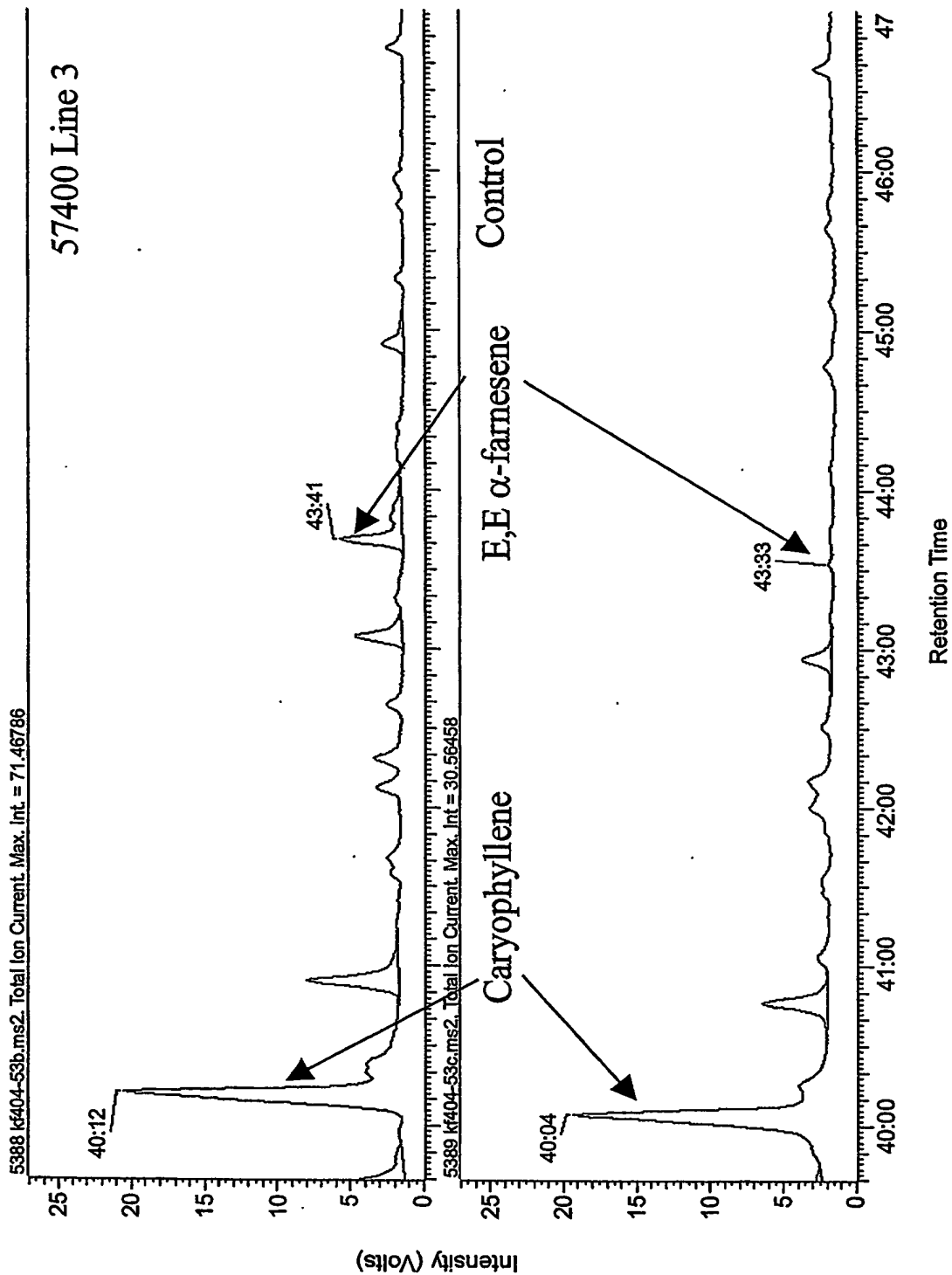


FIGURE 19

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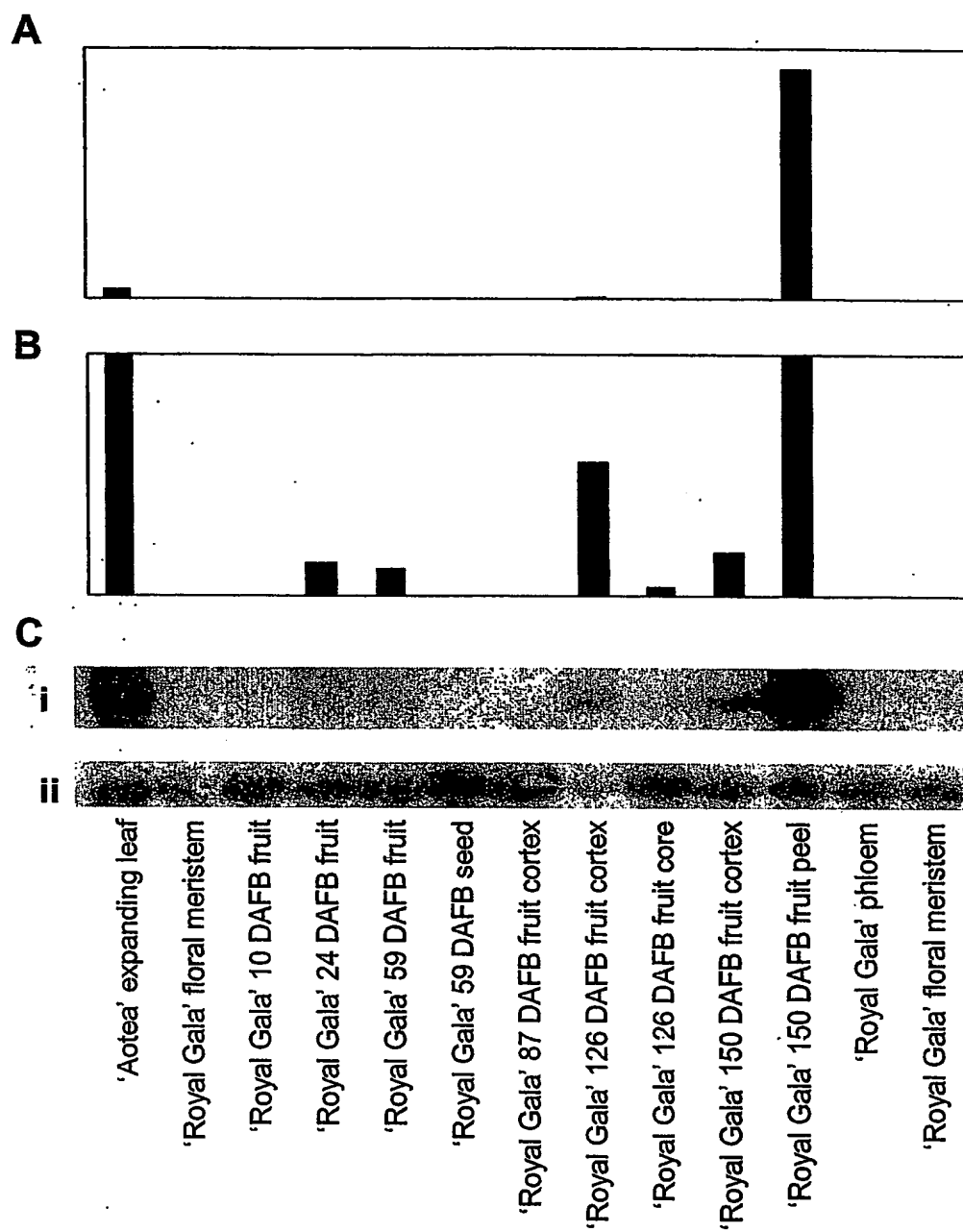


FIGURE 20

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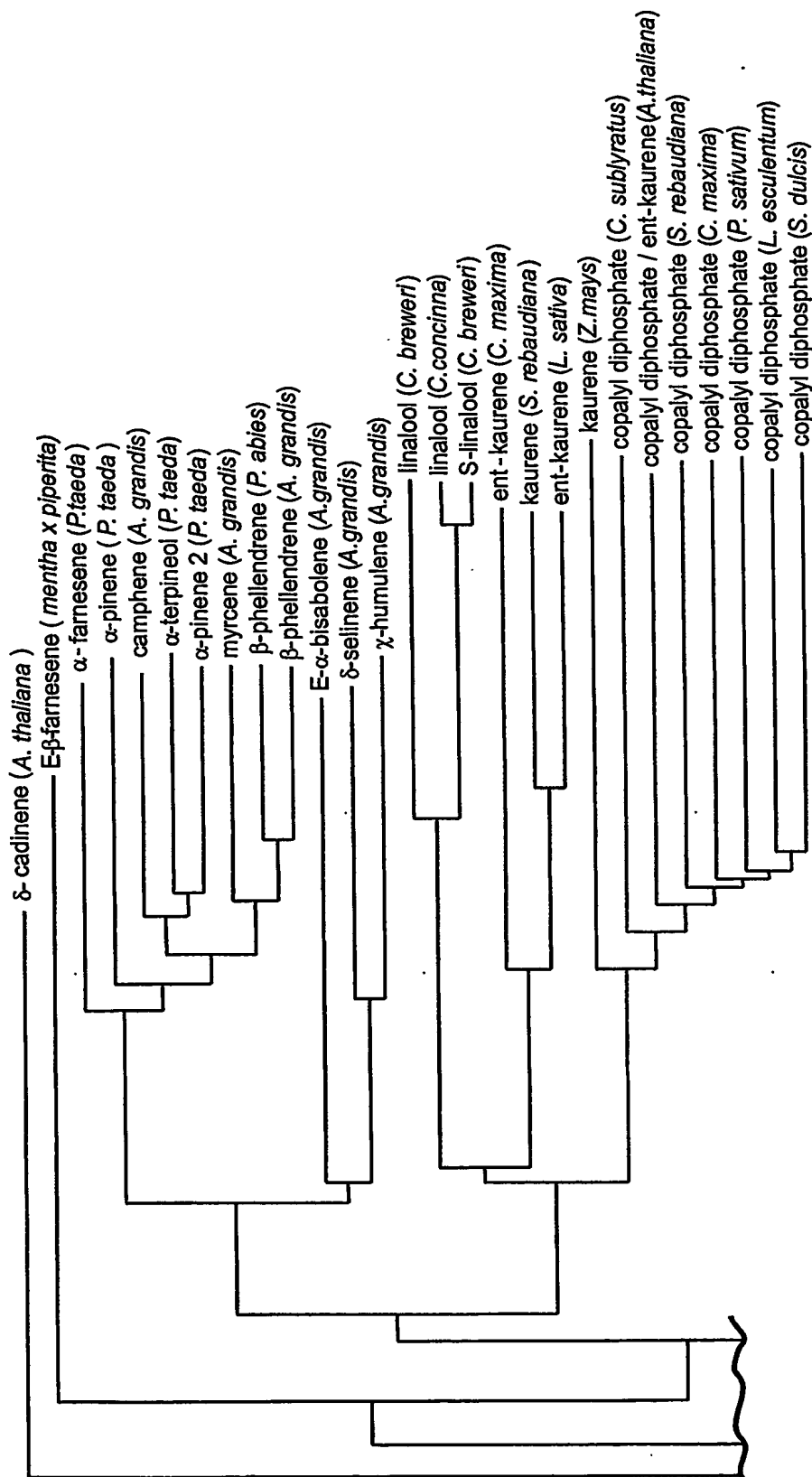


FIGURE 21 (continued next page)

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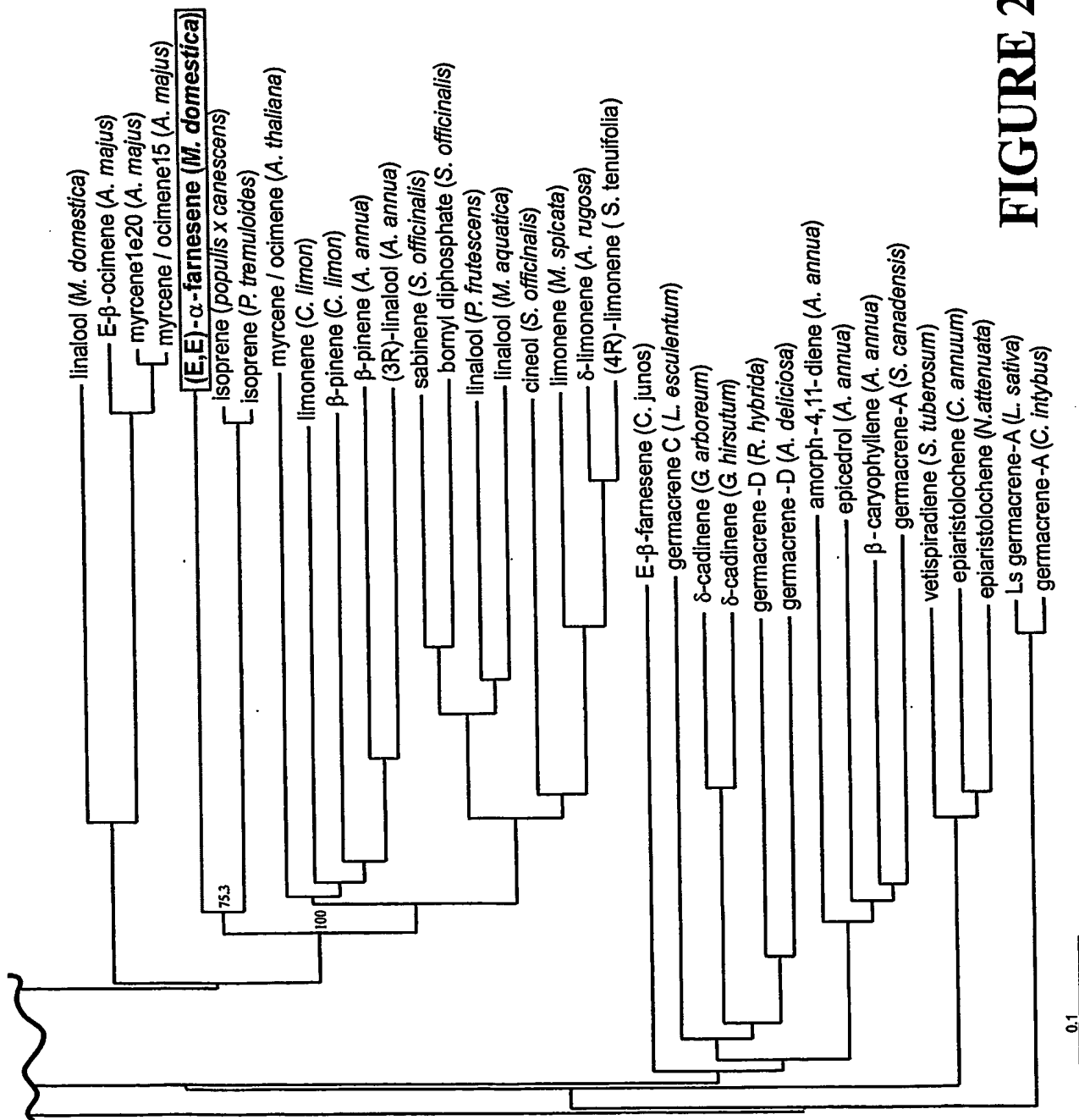


FIGURE 21 (continuation)